

Figure 1

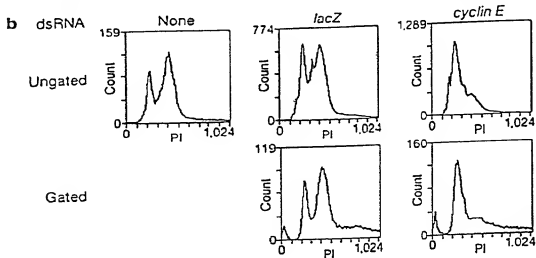
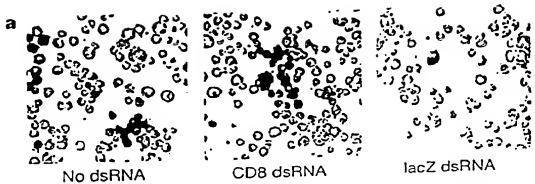


Figure 2

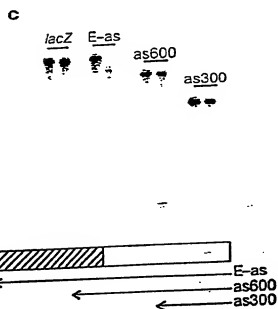
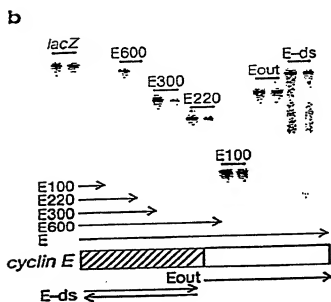
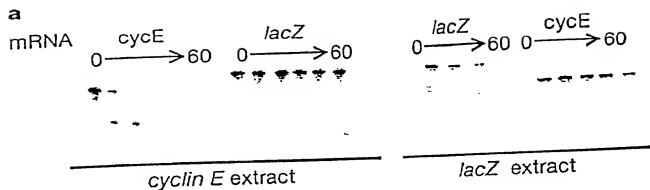


Figure 3

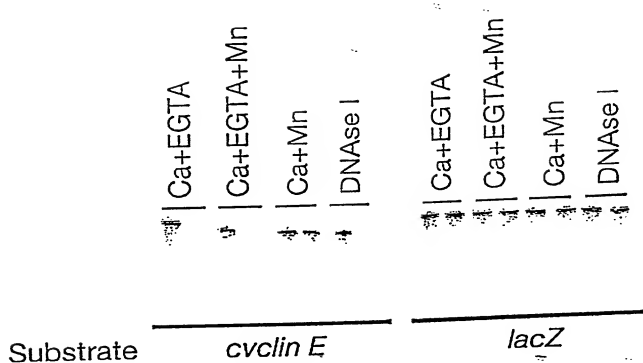
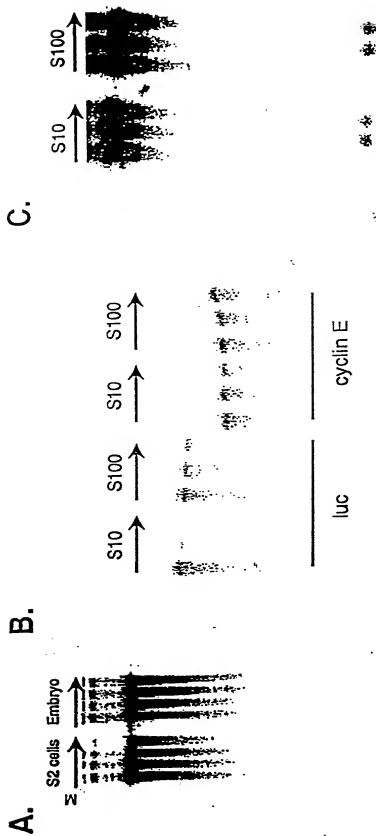
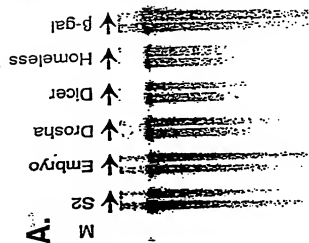


Figure 1 illustrates the steps of the proposed algorithm for finding a minimum spanning tree. The process starts with an initial graph (a) and proceeds through a series of edge selections (b-l) based on the minimum weight rule, resulting in a final minimum spanning tree (l).

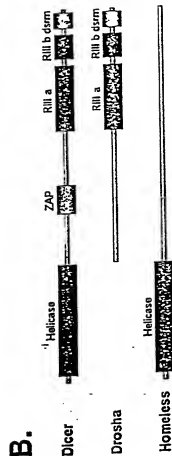


Figure 5





marker
pre-immune
immune
plus peptide
extract



Dicer IP
RISC
control
marker

F.

RISC - hs
RISC - ls

total

E.

D. IP Ext
ATP - + - +

Figure 6d-f

Figure 7

A. casp9 dsRNA
dicer dsRNA

B.

casp9 dsRNA
dicer dsRNA

C.

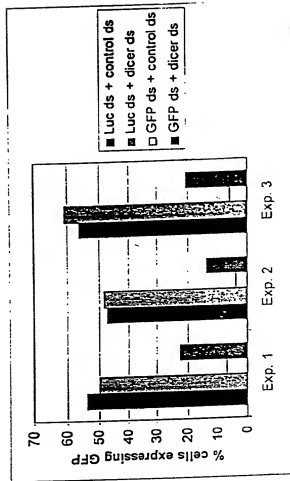


Figure 9

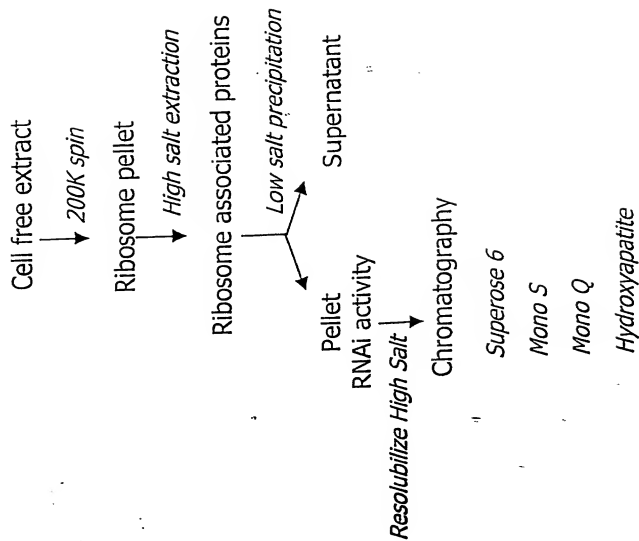


Figure 10

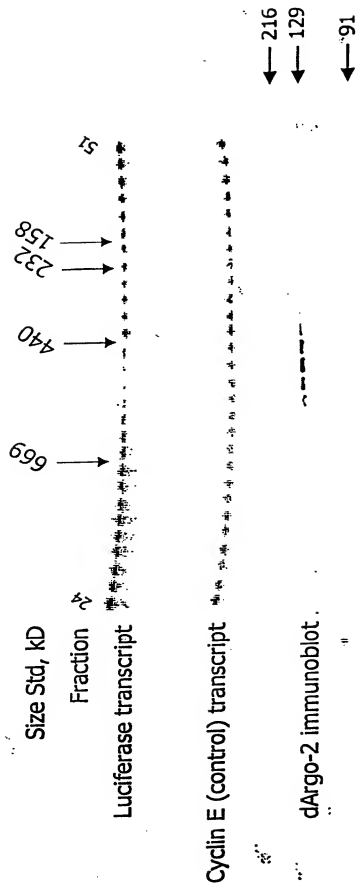


Figure 11

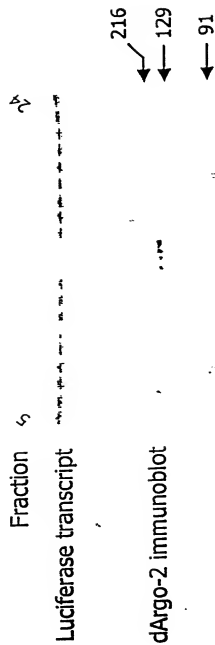


Figure 12

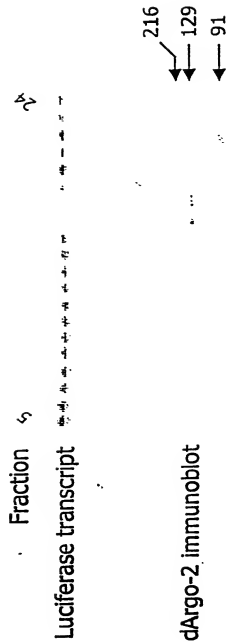


Figure 13

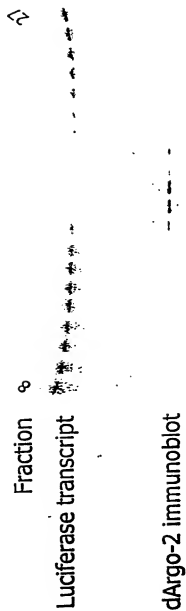
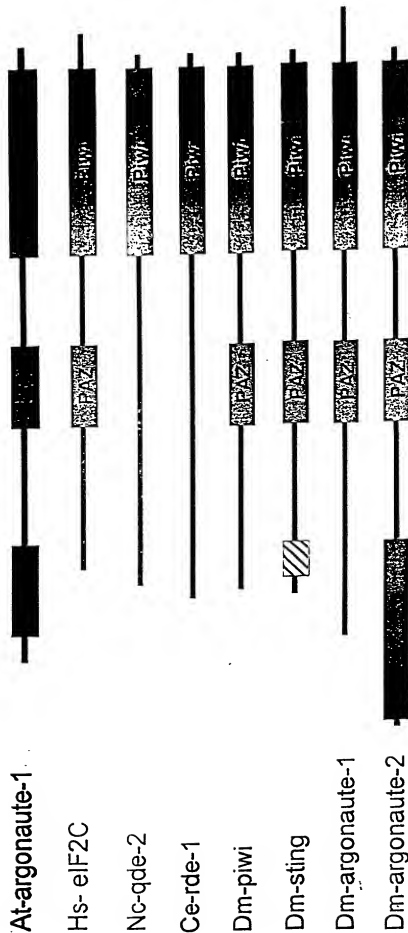


Figure 14



202210-26255001

Figure 15

Ago - high salt
Ago - low salt

total

17

Figure 16

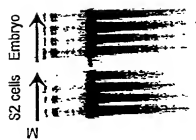


Figure 17

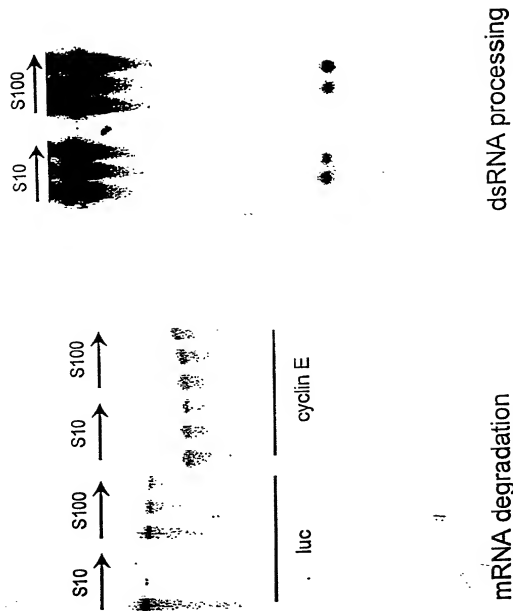
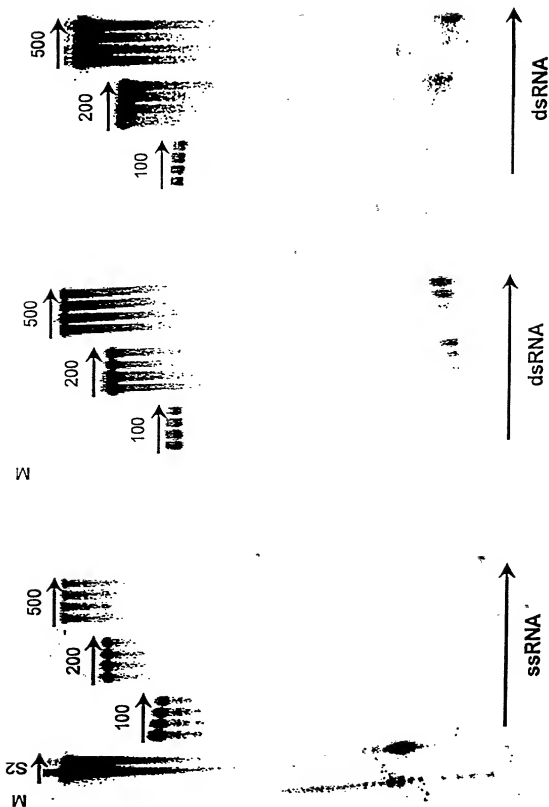


Figure 18

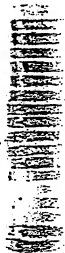


202210-26255001

Resource Phenyl

Q-sepharose

HAP



Superose

S-sepharose

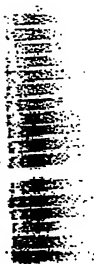
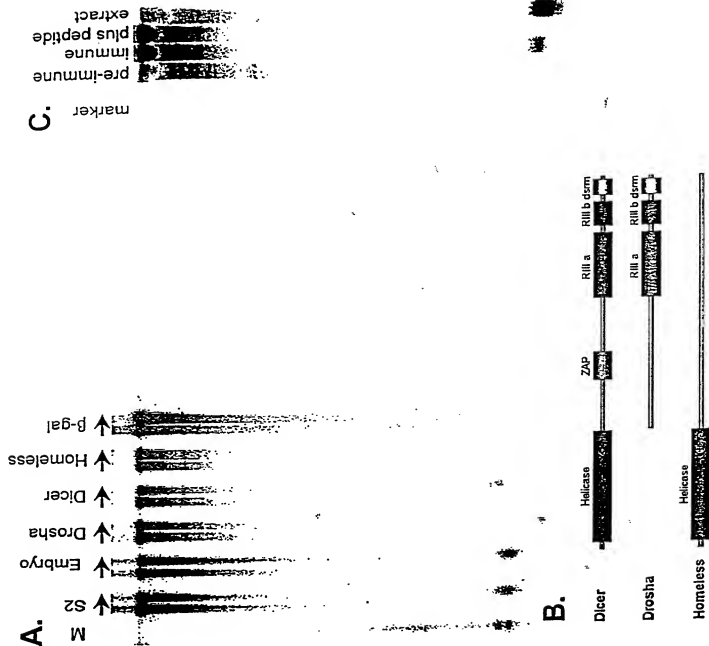


Figure 19

Purification of the 22-mer generating enzyme

Figure 20



202210 1625501

Figure 21

D.

	IP	Ext
ATP	+	+



Figure 22

Dicer IP
RISC
control
marker

1

RISC - hs
RISC - ls

total

1

Figure 23

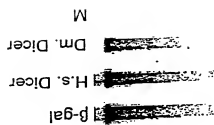


Figure 24

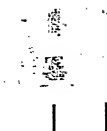
MGKKDKKGGDSAAAPQ000K000R00Q00LQ0PQ0LQ0PQ0LQ0PQ0LQ0PQ00000
 QPHQ0000SSR00PSTSSGSRASGFQ0GG00K0SQDAEGWTAQKKQKQ00VQGWTKQ
 G00GGHQ0GRG0DGGYQ0RPPG0Q0GGHQ0GRG0EGGYQ0RPPG0Q0GGHQ0GRG
 QEGGYQ0RPSG0GGHQ0GRG0QEGGYQ0RPPG0Q0GGHQ0GRG0EGGYQ0RPSGQ
 Q0GGHQ0GRG0EGGYQ0RPSG0Q0GGHQ0GRG0QEGGYQ0RPSG0Q0GGHQ0GRGQ
 EGGYQ0RPPG0Q0PNTQ0SQ0GY0SRGPP0Q0QAAPLPLPQ0PAGSIRGTTIGKPGQVG
 INYLDLDSKMP5VAHYDVKIMPERPKFYQAFQ0FRVDOLGGAVLAYDGKASCYS
 VDKLPLNSQNP5EVTDRNGRTLRYTIEIKETGSDTDLKSLTTYMNDRI**FDKPKRAM**
 QCV5VLASCHNKAIRVGRSEFKWSPNNRHELD0GYEALVGLYQAFMLGDRPFLNV
 DISHKSPTISMPMIEYLERSLKAKINNTWLDYSRRLEPFLRGINVVVTPPQSFQS
 APRVYRVNGLSRAPASSETFEHDKGKVTIASYFHSRNYPLKFPQLHCLNVGSSIKSIL
 LPIELCSIEGQALNRKGATQVANMIKYAAT5INVRRKIMNLLQYFQHNLDP5ISR
 FGIRIANDFI5V5TRV5LSPQVEYH5KRFT5VKN5G5WRMDGNK5FLEPKPAHKCAVLY
 CDPR5GRKMNYTQ5LND5GNLII5QCKAVNISLDS0V5YR5P5TDD5ERSL0TII5ADLKR5
 QHDLA5I5I5P0FRISYD5TI5KQKAE5LQHG5LTQ5IKQ5FTVERKCNQ5TI5NILLKINSK
 L5NGINHKIKD0PRLPM5MKN5TWI5GAD5VTHP5SPDQREI5PSV5GV5AASHD5PYGASYNMQY
 RLQ5GAL5EIEDMF5SIT5LEHLRVY5KEYRNAY5PDHII5Y5RDGVS5DQGF5PKIKNEELRCI
 KOACDKVGCKPKICCVI5V5V5KRH5TR5FP5SGD5V5T5SNK5NN5DP5CTV5VDRT5IVHP5NEMQ
 F5FW5SHQAIQGTAKP5TR5NV5IENTGNL5DIDL5QL5TL5YN5LCH5MF5PR5N5R5SV5Y5PAPAYL
 AHL5VAARG5V5Y5L5GT5NR5FL5DLK5EYAKRTI5V5PE5FMKKN5PMYFV

Figure 25



S2 genomic
S2 cDNA
Library clone #7
Argo-2/p12
No template

Embryo
Adult
S2



105577 01202
202207 262500

untransfected
hdicer transfected
Embryo extract

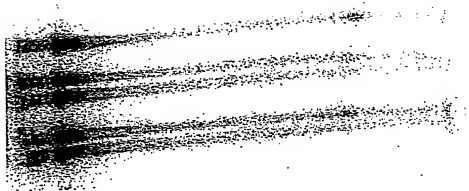
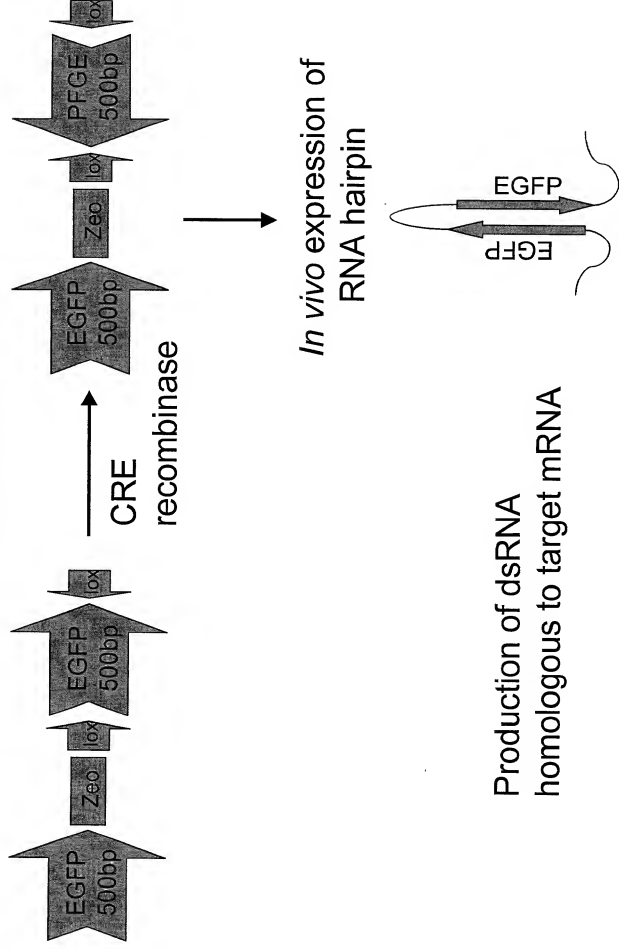


Figure 26

Figure 27

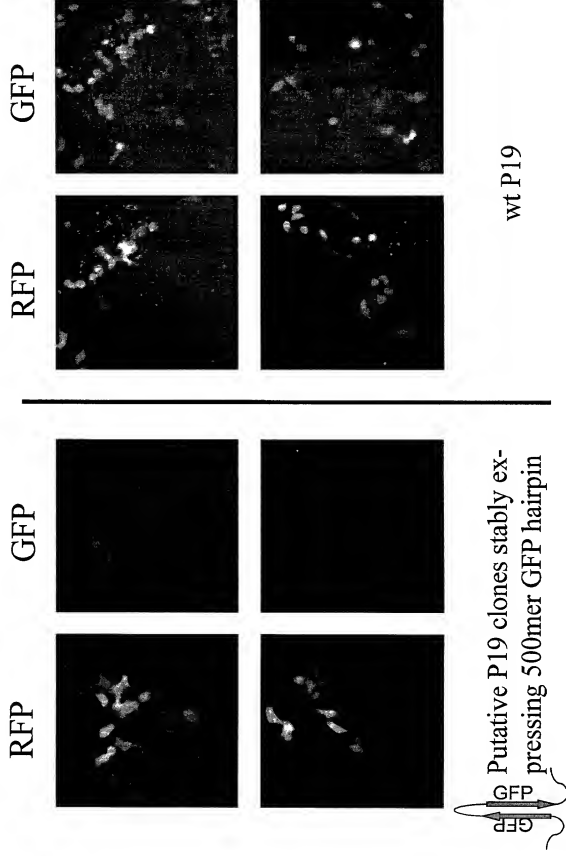
Strategy for stable expression of dsRNA in cultured mammalian cells



Production of dsRNA homologous to target mRNA

Figure 28

Stable suppression of transgene expression in mammalian cells



Co-transfection with pRFP and pGFP, 42 hrs post-transfection

Figure 29

202210-2625501

Dual luciferase assay 21 hrs post-transfection (.4ug dsRNA)

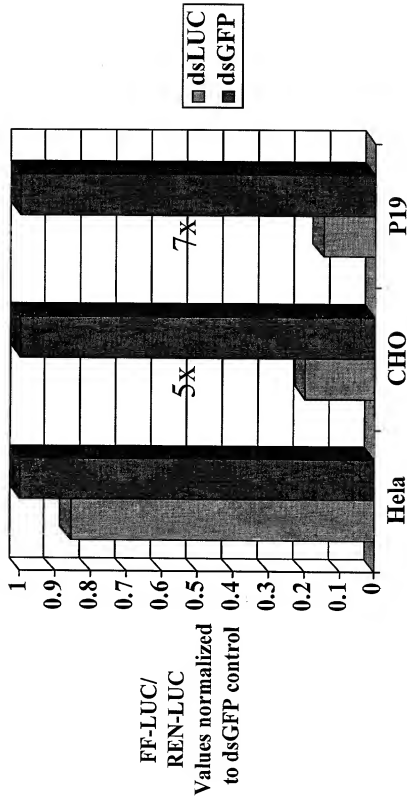


Figure 30
2022102625501
RNAi in ES cells

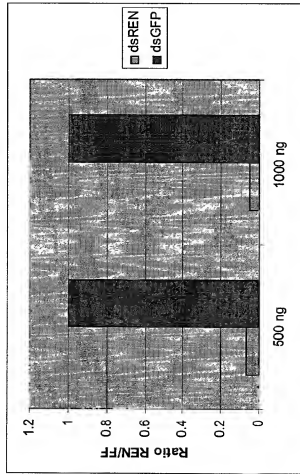


Figure 31

202210-6699501

RNAi in mouse embryonic cells (P19)

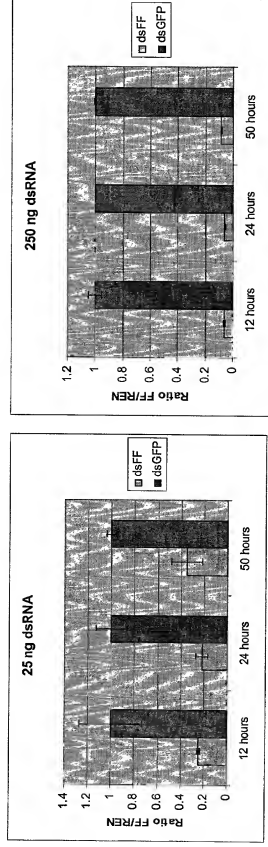


Figure 32

RNAi is post-transcriptional

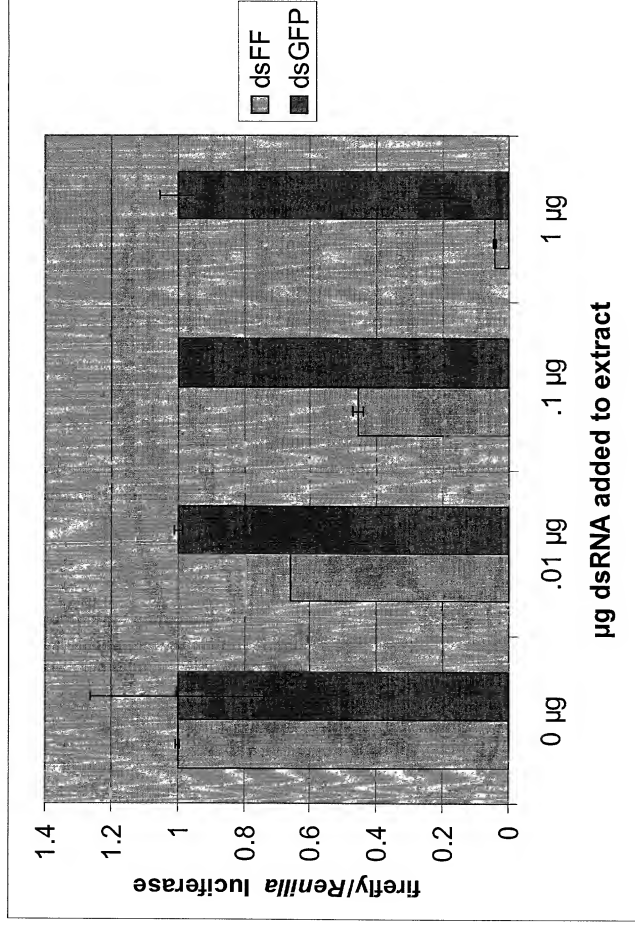
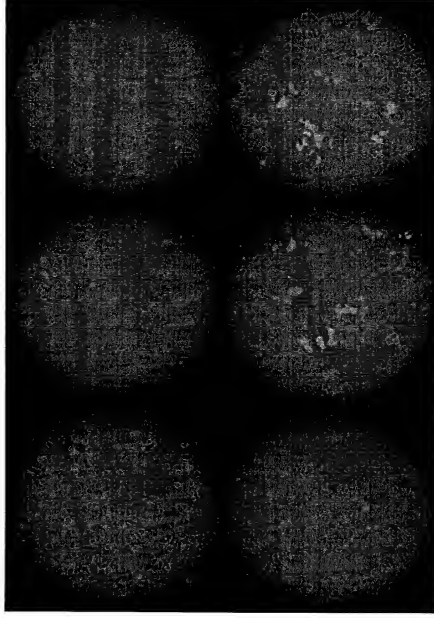


Figure 33

pGFP +	pGFP +	pGFP +
no dsRNA	500ng dsRNA	1000ng dsRNA



dsFF

dsDicer

P19 GFP hairpin clone number #10
48hrs post-transfection
Fluorescent microscopy superimposed with bright field

Figure 34

2022102625501

Silencing is specific and requires dsRNA

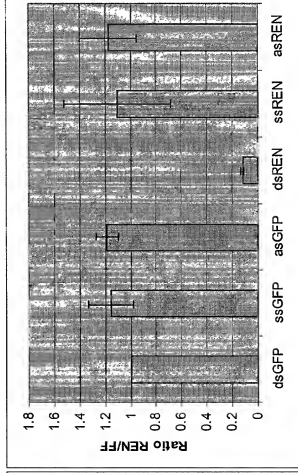
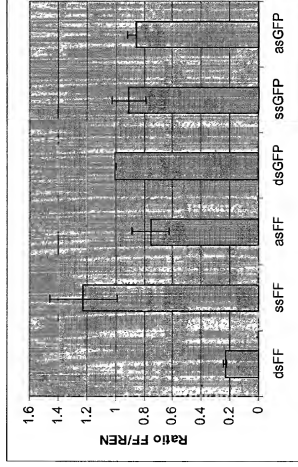


Figure 35

202210-2625001

P19 cells soaked with in dsRNA for
12 hrs in 2mL growth medium (alpha MEM, 10% FBS)

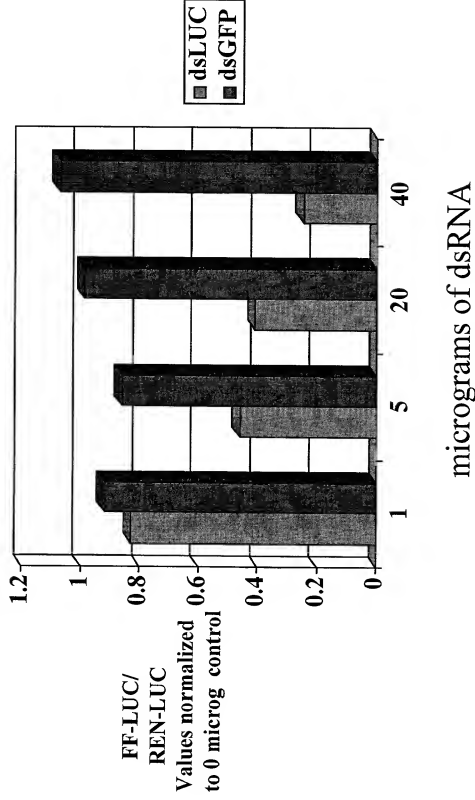
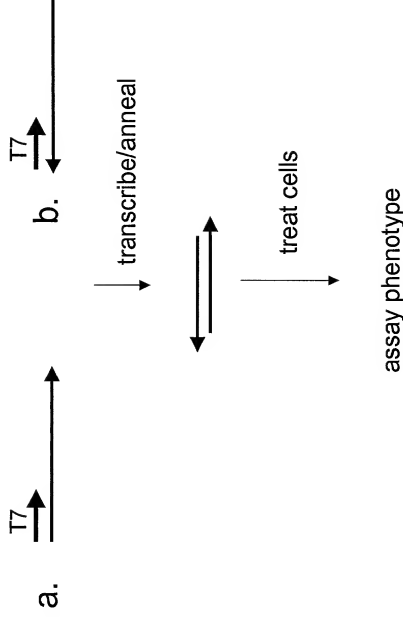


Figure 36

202210-2655001

In vitro synthesis of siRNAs by T7 RNA polymerase



DNA synthesis/RNA transcription

~ \$16/siRNA versus ~\$400/siRNA for chemical synthesis

Brings large-scale projects within reasonable budget range

Figure 37

202210-26ZSSDT

Luciferase siRNA

UUCAGAGCUCAGCGUAGUGA
UAAGGCUACAGGCGAUUC

Luciferase Let-7 like

CAAGAGCGAAUCCUUGGAAUCCGUU
UUAUCCGUUUUAGCGCAUUAUAGUAAA
UAGGUAUUG
U
GGGC
UCCG C
U

Luciferase simple hairpin

U
CAUGAGCGAAUCCUUGGAAUCCGUU
UAGGUAUUGGAAUUAUAGUAAA
A

Figure 38

202410-26Z55301

Short Hairpin RNAs in Drosophila S2 cells

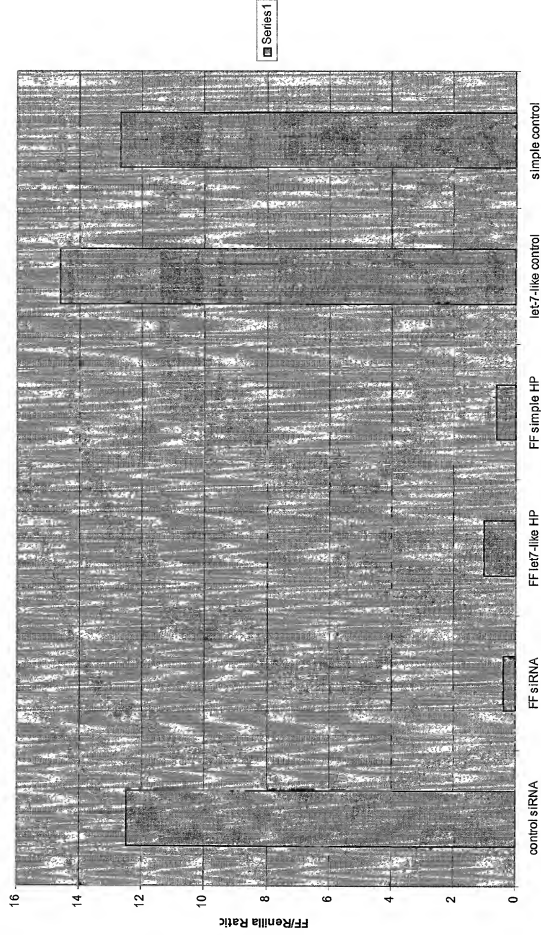


Figure 39

202210-26/55001

Short Hairpin RNAs in Human 293T cells

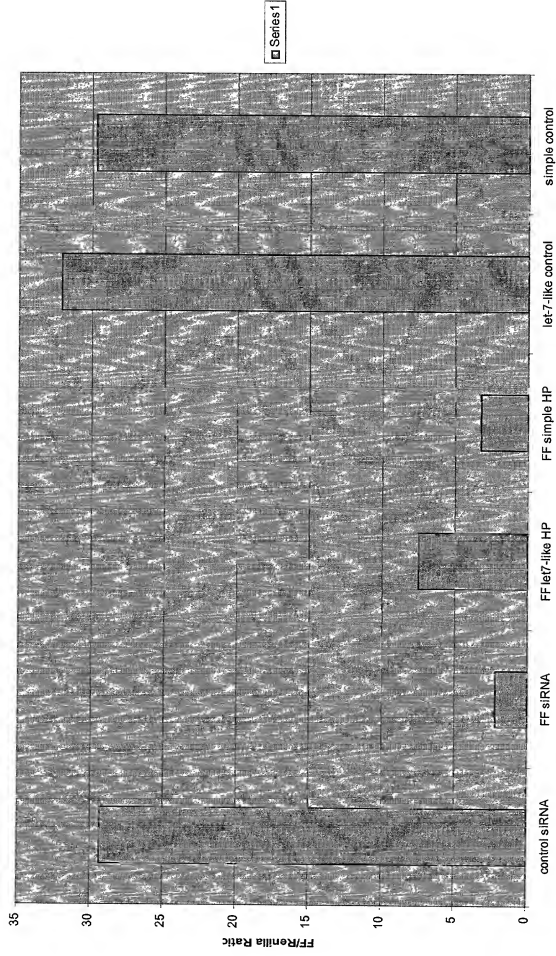


Figure 40

20221026Z55001

Short Hairpin RNAs in Human HeLa cells

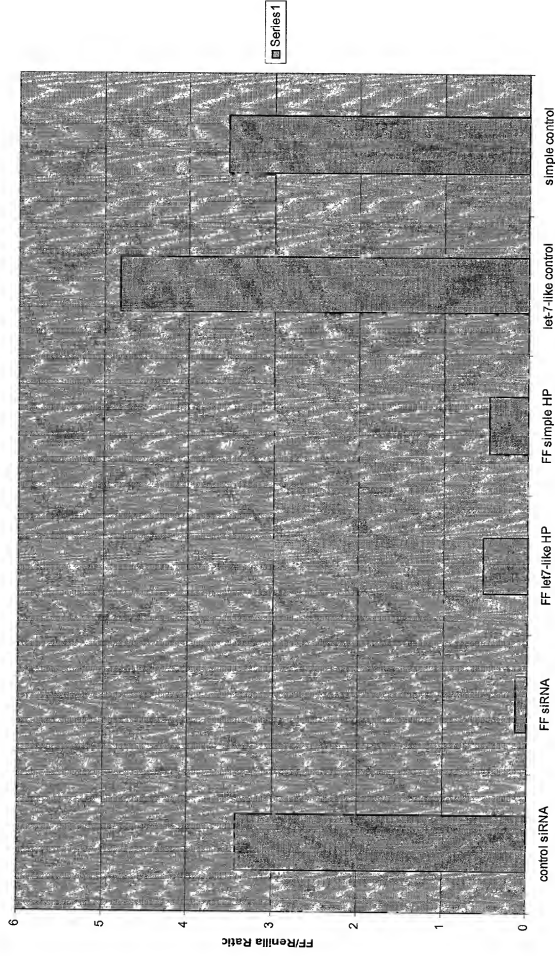


Figure 41

202210*26255001

Simultaneous introduction of multiple hairpins does not produce synergy

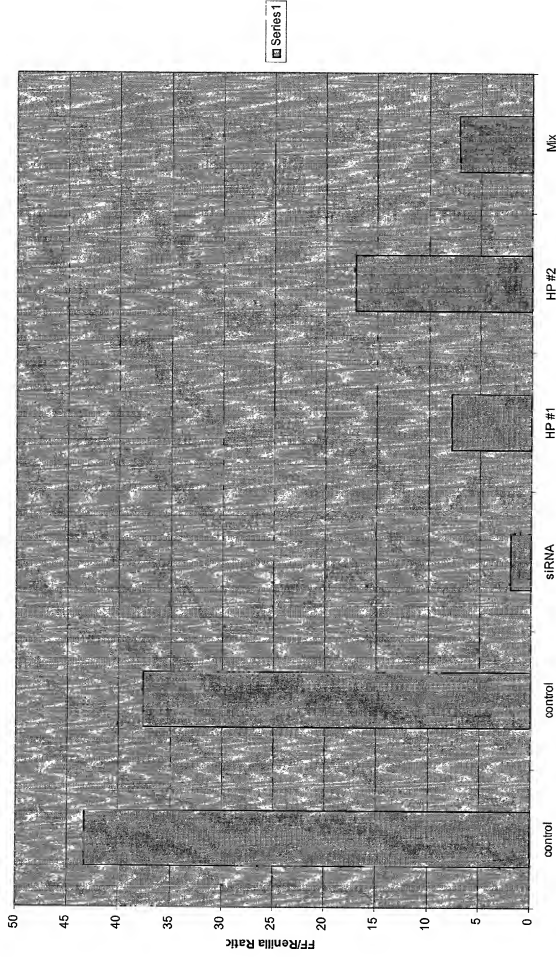


Figure 42
 Encoded short hairpins function *in vivo*

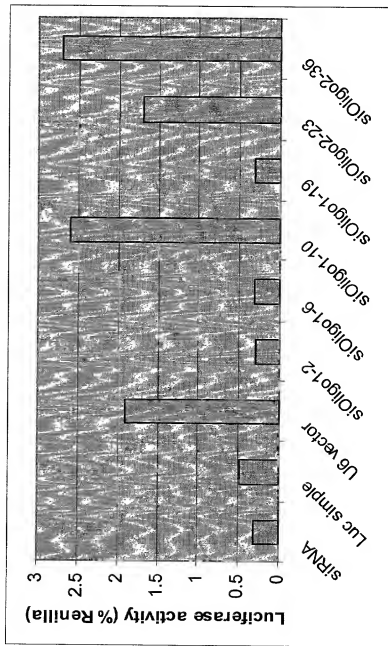
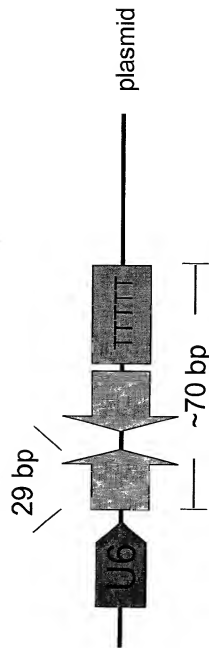


Figure 43

202310-26255001

Stable Suppression by short dsRNAs — stable expression strategies



T7 gives site-specific initiation. 3' end formation
Achieved with ribozyme (e.g. hepatitis delta virus ribozyme).



polII gives site-specific initiation.
Example promoters – U6 snRNA, H1 RNA, SRP RNAs (7SL)
3' end formation
Achieved with native terminator (e.g. TTITT). Leaves the last
TT, so that could be used to pair to transcript.

Could also use VAI, tRNA etc but would have to couple with
Ribozyme since those promoters need also internal elements.

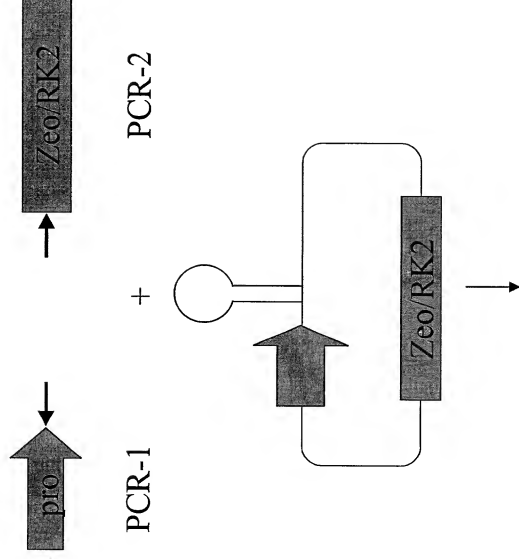


polII gives site-specific initiation. Example promoters
Would be U1 snRNA promoters, CMV etc...
3' end formation achieved with ribozyme
(e.g. hepatitis delta virus ribozyme).

Figure 44

202210-26255001

Stable Suppression by short dsRNAs – cloning strategy

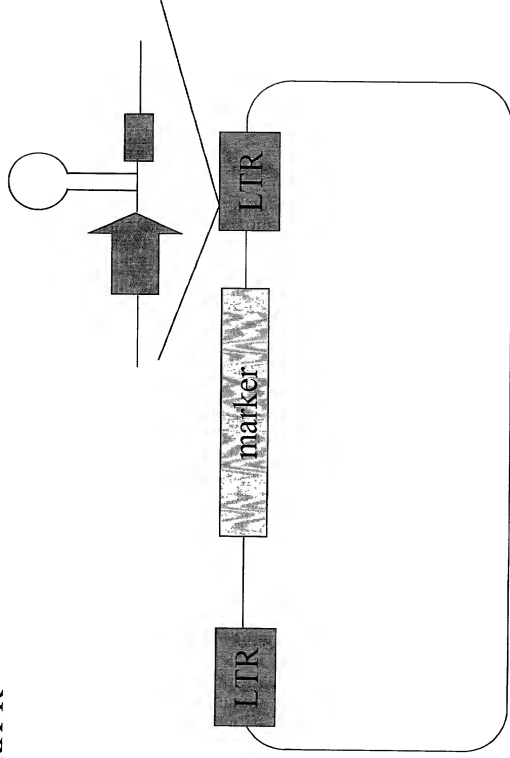


Automatic subcloning into vector of choice

Figure 45

202210-26/55001

MaRX-R



Stable suppression by expressed RNAi

Figure 46

20221026255001

Early Passage PKR^{-/-} MEFs: dual luciferase assay with long dsRNA (~500nt)

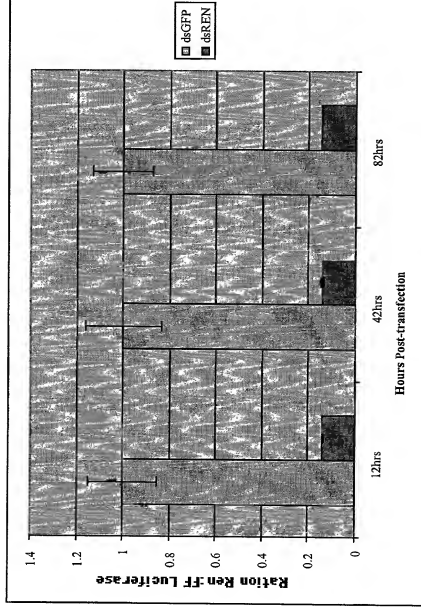


Figure 47

20221016255001

Mouse Tyrosinase Promoter

